AMENDMENTS TO THE CLAIMS

- 1. (currently amended) A chimeric protein comprising:
 - (a) a Kunitz-type domain 1 of TFPI-2 or a mutein thereof and (b) a Kunitz-type domain 2 of TFPI or a mutein thereof; or
 - (c) a Kunitz-type domain 2 of TFPI-2 or a mutein thereof and (d) a Kunitz-type domain 1 of TFPI or a mutein thereof,

wherein the chimeric protein binds and inhibits factor VIIa / tissue factor complex and binds to and inhibits factor Xa.

2. (currently amended) The chimeric protein of claim 1, wherein said chimeric protein is represented by the generic structure:

$$A-(X_1)_a - B-(X_2)_b-C$$

wherein A and C are independently optional flanking peptides, the flanking peptides containing 0-100 amino acids;

wherein B is an optional spacer peptide, the spacer peptide containing 0-25 amino acids; wherein each X_1 is -D- K_1 -E-

where D, E are independently peptides of 0-25 amino acids,

where K_1 comprises TFPI Kunitz-type domain 1 or a mutein thereof, or TFPI-2 Kunitz-type domain 1 or a mutein thereof;

wherein each X2 is -F-K2-G-

where F, G are independently peptides of 0-25 amino acids,

where K₂ comprises TFPI Kunitz-type domain 2 or a mutein thereof, or TFPI-2 Kunitz-type domain 2 or a mutein thereof;

wherein a, b are integers from 1-6 0-6;

wherein A, B, C, D, E, F, G may comprise portions of native TFPI or TFPI 2 sequences or non-native sequence; and

the chimeric protein is not native TFPI or TFPI-2.

- 3. (original) The chimeric protein of claim 2, wherein A or C comprises Kunitz-type domain 3 of TFPI.
- 4. (original) The chimeric protein of claim 2, wherein A or C comprises Kunitz-type domain 3 of TFPI-2.
- 5. (original) The chimeric protein of claim 2, wherein at least one of said flanking peptides comprises an amino acid sequence capable of binding one or more cell surface components.
- 6. (original) The chimeric protein of claim 5, wherein said amino acid sequence capable of binding one more cell surface components is an amino acid sequence capable of binding a glycosaminoglycan.
- 7. (original) The chimeric protein of claim 6, wherein said amino acid sequence capable of binding a glycosaminoglycan is an amino acid sequence capable of binding heparin.
- 8. (original) The chimeric protein of claim 7, wherein said amino acid sequence capable of binding heparin is a heparin-binding domain from a protein, said protein selected from the group consisting of:
 - (a) protease nexin-1;
 - (b) protease nexin-2;
 - (c) antithrombin III;
 - (d) heparin cofactor II;

		(e) protein C inhibitor;
		(f) platelet factor 4;
		(g) bovine pancreatic trypsin inhibitor; and
		(h) ghilanten-related inhibitors.
9.	(original)	The chimeric protein of claim 7, wherein said amino acid sequence capable of
bir	nding hepar	in is a heparin-binding domain selected from the group consisting of:
		(a) SEQ ID NO: 10;
		(b) SEQ ID NO: 11;
		(c) SEQ ID NO: 12;
		(d) SEQ ID NO: 13;
		(e) SEQ ID NO: 14;
		(f) SEQ ID NO: 15;
		(g) SEQ ID NO: 16;
		(h) SEQ ID NO: 17; and
		(i) SEQ ID NO: 18.
10	. (original)) The chimeric protein of claim 5, wherein said flanking peptide comprises the C-
tei	rminal tail o	of TFPI [SEQ ID NO: 7].
11	. (original) The chimeric protein of claim 5, wherein said flanking peptide comprises the C-
tei	rminal tail o	of TFPI-2 [SEQ ID NO: 8].
12	2. (canceled	i)

- 13. (currently amended) The chimeric protein of claim 2, wherein each K_1 is mutein of Kunitz-type domain 1 of TFPI-2, and each K_2 is a mutein of Kunitz-type domain 2 of TFPI, and a and b are integers greater than 1.
- 14. (previously amended) A chimeric protein, wherein the primary amino acid sequence of the chimeric protein is SEQ ID NO: 19.
- 15. (previously amended) The chimeric protein of claim 14, wherein the chimeric protein comprises first and second amino acid sequences, said first amino acid sequence comprising SEQ ID NO:19 and said second amino acid sequence selected from the group consisting of:
 - (a) SEQ ID NO: 7;
 - (b) SEQ ID NO: 8;
 - (c) SEQ ID NO: 10;
 - (d) SEQ ID NO: 11;
 - (e) SEQ ID NO: 12;
 - (f) SEQ ID NO: 13;
 - (g) SEQ ID NO: 14;
 - (h) SEQ ID NO: 15;
 - (i) SEQ ID NO: 16;
 - (i) SEQ ID NO: 17; and
 - (k) SEQ ID NO: 18.

16. (currently amended) The chimeric protein of claim 1, wherein said chimeric protein is represented by the generic structure:

$$A - [X_1 - B - X_2]_e - C$$
 $A - (X_1 - B - X_2)_c - C$

wherein A and C are independently optional flanking peptides, the flanking peptides containing 1-100 amino acids;

wherein B is an optional spacer peptide, the spacer peptide containing 1-25 amino acids; wherein each X_1 is -D- K_1 -E-

where D, E are independently peptides of 1-25 amino acids,

where K_1 is (a) the Kunitz-type domain 1 of TFPI-2 or the mutein thereof or (b) the TFPI Kunitz-type domain 1 of TFPI or the mutein thereof;

wherein each X₂ is -F-K₂-G-

where F, G are independently peptides of 1-25 amino acids,

where K_2 is (a) the Kunitz-type domain 2 of TFPI or the mutein thereof or (b) the Kunitz-type domain 2 of TFPI-2 or the mutein thereof,

wherein c is an integer from 1-10.

- 17. (original) The chimeric protein of claim 16, wherein A or C comprises Kunitz-type domain 3 of TFPI [SEQ ID NO: 7].
- 18. (original) The chimeric protein of claim 16, wherein A or C comprises Kunitz-type domain 3 of TFPI-2 [SEQ ID NO: 8].
- 19. (original) The chimeric protein of claim 16, wherein at least one of said flanking peptides comprises an amino acid sequence capable of binding one or more cell surface components.

- 20. (original) The chimeric protein of claim 19, wherein said amino acid sequence capable of binding one or more cell surface components is an amino acid sequence that binds glycosaminoglycan.
- 21. (original) The chimeric protein of claim 20, wherein said amino acid sequence capable of binding glycosaminoglycan is an amino acid sequence capable of binding heparin.
- 22. (original) The chimeric protein of claim 21, wherein said amino acid sequence capable of binding heparin is a heparin-binding domain from a protein, said protein selected from the group consisting of:
 - (a) protease nexin-1;
 - (b) protease nexin-2;
 - (c) antithrombin III;
 - (d) heparin cofactor II;
 - (e) protein C inhibitor;
 - (f) platelet factor 4;
 - (g) bovine pancreatic trypsin inhibitor; and
 - (h) ghilanten-related inhibitors.
- 23. (original) The chimeric protein of claim 21, wherein said amino acid sequence capable of binding heparin is a heparin-binding domain selected from the group consisting of:
 - (a) SEQ ID NO: 10;
 - (b) SEQ ID NO: 11;
 - (c) SEQ ID NO: 12;
 - (d) SEQ ID NO: 13;

- (e) SEQ ID NO: 14;
- (f) SEQ ID NO: 15;
- (g) SEQ ID NO: 16;
- (h) SEQ ID NO: 17; and
- (i) SEQ ID NO: 18.
- 24. (original) The chimeric protein of claim 19, wherein said flanking peptide comprises the C-terminal tail of TFPI [SEQ ID NO: 7].
- 25. (original) The chimeric protein of claim 19, wherein said flanking peptide comprises the C-terminal tail of TFPI-2 [SEQ ID NO: 8].
- 26. (original) The chimeric protein of claim 1 wherein said protein is produced in a yeast cell and contains no carbohydrate which is immunogenic in mammals.
- 27. (original) The chimeric protein of claim 26 wherein said protein contains no α -1,6-polymannose terminal carbohydrate.
- 28-72. (canceled)
- 73. (original) A pharmaceutical composition comprising the chimeric protein of claim 1 and a pharmaceutically acceptable carrier.
- 74-87. (canceled)
- 88. (previously added) The chimeric protein of claim 2 wherein each K₁ is a mutein of Kunitz-type domain 1 of TFPI and each K₂ is a mutein of Kunitz-type domain 2 of TFPI-2.